

Gambel

CRF Error Corrected by the STIC Systems Branch

Serial Number: 09/027,205A

CRF Processing Date: 7/21/99
Edited by: [signature]
Verified by: [signature] (STIC staff)

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: 18 6/12
- ☐ Edited the Current Application Data section with the actual current number. The number input by the applicant was ☐ the prior application data; or ☐ other ENTERED
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: _____
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: _____
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: _____
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: _____
- ☐ Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file;
☐ page numbers throughout text; ☐ other invalid text, such as _____
- ☐ Inserted mandatory headings, specifically: _____
- ☒ Corrected an obvious error in the response, specifically:
Seq 7 - corrected (A) LENGTH: response
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: _____
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted **ending** stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____
- ☐ Other: _____

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95

RAW SEQUENCE LISTING
PATENT APPLICATION US/09/027,205A

DATE: 07/21/1999
TIME: 14:48:45

INPUT SET: S32598.raw

This Raw Listing contains the General
Information Section and those Sequences
containing ERRORS.

SEQUENCE LISTING

Does Not Comply
Corrected Diskette Needed

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- (1) General Information:
 - (i) APPLICANT: Carl H. June et al.
 - (ii) TITLE OF INVENTION: Methods for Modulating Expression of an HIV-1 Fusion Cofactor
 - (iii) NUMBER OF SEQUENCES: 7
 - (iv) CORRESPONDENCE ADDRESS:
 - (A) ADDRESSEE: LAHIVE & COCKFIELD, LLP
 - (B) STREET: 28 State Street
 - (C) CITY: Boston
 - (D) STATE: Massachusetts
 - (E) COUNTRY: USA
 - (F) ZIP: 02109
 - (v) COMPUTER READABLE FORM:
 - (A) MEDIUM TYPE: Floppy disk
 - (B) COMPUTER: IBM PC compatible
 - (C) OPERATING SYSTEM: PC-DOS/MS-DOS
 - (D) SOFTWARE: PatentIn Release #1.0, Version #1.25
 - (vi) CURRENT APPLICATION DATA:
 - (A) APPLICATION NUMBER:
 - (B) FILING DATE:
 - (C) CLASSIFICATION:
 - (vii) PRIOR APPLICATION DATA:
 - (A) APPLICATION NUMBER: US 60/037,422
 - (B) FILING DATE: 21-FEB-1997
 - (viii) ATTORNEY/AGENT INFORMATION:
 - (A) NAME: Amy E. Mandragouras
 - (B) REGISTRATION NUMBER: 36,207
 - (C) REFERENCE/DOCKET NUMBER: GIN-005
 - (ix) TELECOMMUNICATION INFORMATION:
 - (A) TELEPHONE: (617)227-7400
 - (B) TELEFAX: (617)742-4214

ERRORED SEQUENCES FOLLOW:

RAW SEQUENCE LISTING
PATENT APPLICATION US/09/027,205ADATE: 07/21/1999
TIME: 14:48:45

INPUT SET: S32598.raw

45 (2) INFORMATION FOR SEQ ID NO:1:

46

47 (i) SEQUENCE CHARACTERISTICS:

48 (A) LENGTH: 7 amino acids

49 (B) TYPE: amino acid

50 (D) TOPOLOGY: linear

51

52 (ii) MOLECULE TYPE: peptide

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54 (v) FRAGMENT TYPE: internal

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57 (ix) FEATURE:

58 (A) NAME/KEY: Peptide

59 (B) LOCATION: 1 - 20

60 (D) OTHER INFORMATION: /note= "any 20 Xaas at location 1 - 20 may be abse

61

62 (ix) FEATURE:

63 (A) NAME/KEY: Peptide

64 (B) LOCATION: 26 - 45

65 (D) OTHER INFORMATION: /note= "any 20 Xaas at location 26 - 45 may be abs

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68 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

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70 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa

71 1 5 10 15

72 Xaa Xaa Xaa Xaa Pro Pro Tyr Tyr Leu Xaa Xaa Xaa Xaa Xaa Xaa Xaa

73 20 25 30

74 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa

75 35 40 45

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SEQUENCE VERIFICATION REPORT
PATENT APPLICATION US/09/027,205A

DATE: 07/21/1999

TIME: 14:48:46

INPUT SET: S32598.raw

Line	Error	Original Text
48	Entered (7) and Calc. Seq. Length (45) differ	(A) LENGTH: 7 amino acids